

<!--StartFragment-->RESULT 3
S51398
hypothetical protein YLR260w - yeast (Saccharomyces cerevisiae)
N;Alternate names: hypothetical protein L8479.7
C;Species: Saccharomyces cerevisiae
C;Date: 05-May-1995 #sequence_revision 12-May-1995 #text_change 09-Jul-2004
C;Accession: S51398
R;Miller, N.
submitted to the EMBL Data Library, November 1994
A;Description: The sequence of S. cerevisiae cosmid 8479.
A;Reference number: S51395
A;Accession: S51398
A;Molecule type: DNA
A;Residues: 1-687 <MIL>
A;Cross-references: UNIPROT:Q06147; UNIPARC:UPI000006977B; EMBL:U17244; NID:g577171; PIDN:
AAB67377.1; PID:g577178; GSPDB:GN00012; MIPS:YLR260w
C;Genetics:
A;Gene: SGD:LCB5; MIPS:YLR260w
A;Cross-references: SGD:S0004250
A;Map position: 12R

Query Match 18.5%; Score 356.5; DB 2; Length 687;
Best Local Similarity 24.9%; Pred. No. 5.6e-24;
Matches 102; Conservative 76; Mismatches 150; Indels 81; Gaps 11;

Qy	1	VLVLLNPRGGKGKALQLFRSHVQPLLAEEAISFTLMLTERRNHARELVRSEELGRWDALV	60
	:	: : : : : :: : : :: :: :	
Db	271	IFVIIINPFGGKGAKKLFMTKAKPLLLASRCSIEVVYTKYPGHAIEIAREMDIDKYDTIA	330
Qy	61	VMSGDGLMHEVVNGLMERPDWETAIQK-PLCSLPAGSGNALAASLNHYAGYEQVTNEDLL	119
	: : : : : : :		
Db	331	CASGDGIPHEVINGLYQRPDHVKAFNNIAITEIPCGSGNAMSVSC-HW-----TNNP--	381
Qy	120	TNCTLLLCCRLLSPMNLLSLHTASGLR---LFSVLSLAWGFIADVLESEKYRRLGEMRF	176
	: : : : :: : : : :		
Db	382	SYSTLCLIKSIETRIDLMCCSQPSYAREHPKLSFLSQTYGLIAETDINTEFIRWMGPARF	441
Qy	177	TLGTFLRLAALRTY-----RGRLAYLPV-----	199
	: : : : : :		
Db	442	ELGVAFNIIQKKKYPCEIYVKYAAKSKNELKNHYLEHKNKGSLEFQHITMNKDNECDNY	501
Qy	200	---GRVGSKTPASPVVVQQGPVDAHLV-----PLEEPVPSHWTW	235
	:: : : : : : : :		
Db	502	NYENEYETENEDEDEDADADDEDSHLISRDLADSSADQIKEEDFKIKYPLDEGIPSDWER	561
Qy	236	VPDEDFVLVLLALLHS---HLGSEMFAAPMGRCAAGVMHLFYVRAGVSRAMLLRLFLAME	291
	: : : :: : : : : : ::		
Db	562	L-DPNISNNLGIFYTGKMPYVAADTKFFPAALPSDGTMDMVITDARTSLTRMAPILLGLD	620
Qy	292	KGRHMEYECPYLVYVPVVAFRLEPKDGKGVFAVDGELMVSEAVQGQVHP	340
	: : :: : : : :		
Db	621	KGSHVQLQ--PEVLHSKILAYKIIPKLGNGLFSVDGEKFPLEPLQVEIMP	667

<!--EndFragment-->